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Title: Haplotypes of the TNFRSF11B Gene

Application No.: To be assigned

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For: Haplotypes of the TNFRSF11B Gene

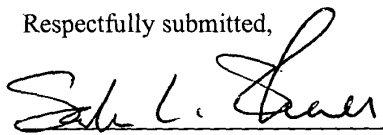
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STATEMENT ACCOMPANYING "SEQUENCE LISTING"  
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The undersigned hereby states upon information and belief that the information recorded in the computer readable form of the sequence listing is identical to the written (paper) sequence listing submitted with the application.

Respectfully submitted,

9-January-2002  
Date

  
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## POLYMORPHISMS IN THE TNFRSF11B GENE

|                 |            |            |             |            |      |
|-----------------|------------|------------|-------------|------------|------|
| ACAGCGAACC      | CTAGAGCAAA | GTGCCAAACT | TCTGTCGATA  | GCTTGAGGCT |      |
| AGTGGAAAGA      | CCTCGAGGAG | GCTACTCCAG | AAGTTCAGCG  | CGTAGGAAGC | 100  |
| TCCGATACCA      | ATAGCCCTTT | GATGATGGTG | GGGTTGGTGA  | AGGGAACAGT |      |
| GCTCCGCAAG      | GTTATCCCTG | CCCAGGCAG  | TCCAATTTTC  | ACTCTGCAGA | 200  |
| TTCTCTCTGG      | CTCTAACTAC | CCCAGATAAC | AAGGAGTGAA  | TGCAGAATAG |      |
| CACGGGCTTT      | AGGGCCAATC | AGACATTAGT | TAGAAAAATT  | CCTACTACAT | 300  |
| GGTTTATGTA      | AACCTGAAGA | TGAATGATTG | CGAACTCCCC  | GAAAAGGGCT |      |
| CAGACAATGC      | CATGCATAAA | GAGGGGCCCT | GTAATTTGAG  | GTTTCAGAAC | 400  |
| CCGAAGTGAA      | GGGGTCAGGC | AGCCGGGTAC | GGCGGAAACT  | CACAGCTTTC |      |
| GCCCAGCGAG      | AGGACAAAGG | TCTGGGACAC | ACTCCAACCTG | CGTCCGGATC | 500  |
| TTGGCTGGAT      | CGGACTCTCA | GGGTGGAGGA | GACACAAGCA  | CAGCAGCTGC |      |
| T               |            |            |             |            |      |
| CCAGCGTGTG      | CCCAGCCCTC | CCACCGCTGG | TCCCGGCTGC  | CAGGAGGCTG | 600  |
| GCCGCTGGCG      | GGAAGGGGCC | GGGAAACCTC | AGAGCCCCGC  | GGAGACAGCA |      |
| GCCGCCTTGT      | TCCTCAGCCC | GGTGGCTTTT | TTTTCCCCTG  | CTCTCCCAGG | 700  |
| GGACAGACAC      | CACCGCCCCA | CCCTCACGC  | CCCACCTCCC  | TGGGGGATCC |      |
|                 | T          |            |             | T          |      |
| TTTCCGCCCC      | AGCCCTGAAA | GCGTTAATCC | TGGAGCTTTC  | TGCACACCCC | 800  |
|                 | C          |            |             |            |      |
| CCGACCGCTC      | CCGCCCAAGC | TTCCTAAAAA | AGAAAGGTGC  | AAAGTTTGGT |      |
| CCAGGATAGA      | AAAATGACTG | ATCAAAGGCA | GGCGATACTT  | CCTGTTGCCG | 900  |
| GGACGCTATA      | TATAACGTGA | TGAGCGCACG | GGCTGCGGAG  | ACGCACCGGA |      |
| GCGCTCGCCC      | AGCCGCCGCC | TCCAAGCCCC | TGAGGTTTCC  | GGGGACCACA | 1000 |
| ATGAACAAGT      | TGCTGTGCTG | CGCGCTCGTG | GTAAGTCCCT  | GGGCCAGCCG |      |
| C               |            |            |             | T          |      |
| [exon 1: 1001.. |            |            |             |            |      |
| ..1030]         |            |            |             |            |      |
| ACGGGTGCCC      | GGCGCCTGGG | GAGGCTGCTG | CCACCTGGTC  | TCCCAACCTC | 1100 |
| CCAGCGGACC      | GGCGGGGAGA | AGGCTCCACT | CGCTCCCTCC  | CAGGAGAGGC |      |
|                 | A          |            |             |            |      |
| TTGGGGTTAG      | GCTGGAGCAG | GAAACCGCTT | TCAAGTTATG  | CCATGCTTCC | 1200 |
| CCTAGGGTGT      | CCTTTTACGC | TGCAAAGTTC | CTGCTGACTT  | TATGGAAGAC |      |
|                 | A          |            |             |            |      |
| AGCAAGAGAG      | AGACAGACAG | CGAGAGAGAG | GGAGAGAGAG  | AGAGAGAGAA | 1300 |
| ACTTGTTTGA      | AAGTTTTAGT | CATTAACCTT | CTGTCTTCAT  | CTCAGAATAT |      |
| TAACGCCCTC      | ATGTAGTCCA | TACTATCTTT | GCTTAATGAA  | CTTGAACCTT | 1400 |
| TATTATTAGT      | GGCAAAGAAG | TGGTCCCTTA | GATTCAGAGT  | AAGTTGGAAG |      |
| AAGACGTTAG      | TCTTCTTAAA | ACCATTATAA | TTAGAATATG  | ACATGATAGA | 1500 |
| NNNNNNNNNN      | NNNNNNNNNN | NNNNNNNNNN | NNNNNNNNNN  | NNNNNNNNNN |      |
| CAGGACTTTG      | AGTCAAATGA | TACTGTTGCA | CATAAGAACA  | AACCTATTTT | 1600 |
| CATGCTAAGA      | TGATGCCACT | GTGTTCCCTT | CTCCTTCTAG  | TTTCTGGACA |      |
| [exon 2: 1641.. |            |            |             |            |      |
| TCTCCATTAA      | GTGGACCACC | CAGGAAACGT | TTCTTCCAAA  | GTACCTTCAT | 1700 |
| TATGACGAAG      | AAACCTCTCA | TCAGCTGTTG | TGTGACAAAT  | GTCCTCCTGG |      |
| TACCTACCTA      | AAACAACACT | GTACAGCAAA | GTGGAAGACC  | GTGTGCGCCC | 1800 |
| CTTGCCCTGA      | CCACTACTAC | ACAGACAGCT | GGCACACCAG  | TGACGAGTGT |      |
| CTATACTGCA      | GCCCCGTGTG | CAAGGAGCTG | CAGTACGTCA  | AGCAGGAGTG | 1900 |
| CAATCGCACC      | CACAACCGCG | TGTGCGAATG | CAAGGAAGGG  | CGCTACCTTG |      |
| AGATAGAGTT      | CTGCTTGAAA | CATAGGAGCT | GCCCTCCTGG  | ATTTGGAGTG | 2000 |
| GTGCAAGCTG      | GTACGTGTCA | ATGTGCAGCA | AAATTAATTA  | GGATCATGCA |      |
|                 | T          |            |             |            |      |
| ..2010]         |            |            |             |            |      |

FIGURE 1A

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10044674.010902

|            |             |            |            |             |      |
|------------|-------------|------------|------------|-------------|------|
| 2/7        |             |            |            |             |      |
| AAGTCAGATA | GTTGTGACAG  | TTTAGGAGAA | CACTTTTGTT | CTGATGACAT  | 2100 |
| TATAGGATAG | CAAATTGCAA  | AGGTAATGAA | ACCTGCCAGG | TAGGTACTAT  |      |
| GTGCTGGAG  | TGCTTCCAAA  | GGACCATTGC | TCAGAGGAAT | ACTTTGCCAC  | 2200 |
| C          |             |            |            |             |      |
| TACAGGGCAA | TTTAATGACA  | AATCTCAAAT | GCAGCAAATT | ATTCTCTCAT  |      |
| GAGATGCATG | ATGGTTTTTT  | TTTTTTTTTT | TAAAGAAACA | AACTCAAGTT  | 2300 |
| GCACTATTGA | TAGTTGATCT  | ATACCTCTAT | ATTTCACTTC | AGCATGGACA  |      |
| CCTTCAAAC  | GCAGCACTTT  | TTGACAAACA | TCAGAAATGT | TAATTTATAC  | 2400 |
| CAAGAGAGTA | ATTATGCTCA  | TATTAATGAG | ACTCTGGAGT | GCTAACAATA  |      |
| AGCAGTTATA | ATTAATTATG  | TAAAAAATGA | GAATGGTGAG | GGGAATTGCA  | 2500 |
| TTTCATTATT | AAAAACAAGG  | CTAGTTCTTC | CTTTAGCATG | GGAGCTGAGT  |      |
| GTTTGGGAGG | GTAAGGACTA  | TAGCAGAATC | TCTTCAATGA | GCTTATTCTT  | 2600 |
| TATCTTAGAC | AAAACAGATT  | GTCAAGCCAA | GAGCAAGCAC | TTGCCTATAA  |      |
| ACCAAGTGCT | TTCTCTTTTG  | CATTTTGAAC | AGCATTGGTC | AGGGCTCATG  | 2700 |
| TGTATTGAAT | CTTTTAAACC  | AGTAACCCAC | GTTTTTTTTC | TGCCACATTT  |      |
| GCGAAGCTTC | AGTGCAGCCT  | ATAACTTTTC | ATAGCTTGAG | AAAATTAAGA  | 2800 |
| GTATCCACTT | ACTTAGATGG  | AAGAAGTAAT | CAGTATAGAT | TCTGATGACT  |      |
| CAGTTTGAAG | CAGTGTCTCT  | CAACTGAAGC | CCTGCTGATA | TTTTAAGAAA  | 2900 |
| TATCTGGATT | CCTAGGCTGG  | ACTCCTTTTT | GTGGGCAGCT | GTCCTGCGCA  |      |
| TTGTAGAATT | TTGGCAGCAC  | CCCTGGACTC | TAGCCACTAG | ATACCAATAG  | 3000 |
| CAGTCCTTCC | CCCATGTGAC  | AGCCAAAAAT | GTCTTCAGAC | ACTGTCAAAT  |      |
| GTCGCCAGGT | GGCAAAATCA  | CTCCTGGTTG | AGAACAGGGT | CATCAATGCT  | 3100 |
| AAGTATCTGT | AACTATTTTA  | ACTCTCAAAA | CTTGTGATAT | ACAAAGTCTA  |      |
| AATTATTAGA | CGACCAATAC  | TTTAGGTTTA | AAGGCATACA | AATGAAACAT  | 3200 |
| TCAAAAATCA | AAATCTATTC  | TGTTTCTCAA | ATAGTGAATC | TTATAAAATT  |      |
| AATCACAGAA | GATGCAAAAT  | GCATCAGAGT | CCCTTAAAT  | TCCTCTTCGT  | 3300 |
| ATGAGTATTT | GAGGGAGGAA  | TTGGTGATAG | TTCTTACTTT | CTATTGGATG  |      |
| GTAATTTGAG | ACTCAAAAGC  | TAAGCTAAGT | TGTGTGTGTG | TCAGGGTGCG  | 3400 |
| GGGTGTGGAA | TCCCATCAGA  | TAAAAGCAAA | TCCATGTAAT | TCATTTCAGTA |      |
| AGTTGTATAT | GTAGAAAAAT  | GAAAAGTGGG | CTATGCAGCT | TGGAACTAG   | 3500 |
| AGAATTTTGA | AAAATAATGG  | AAATCACAA  | GATCTTTCTT | AAATAAGTAA  |      |
| GAAAATCTGT | TTGTAGAATG  | AAGCAAGCAG | GCAGCCAGAA | GAATCAGAAC  | 3600 |
| AAAAGTACAC | ATTTTACTCT  | GTGTACACTG | GCAGCACAGT | GGGATTTATT  |      |
| TACCTCTCCC | TCCCTAAAAA  | CCCACACAGC | GGTTCTCTCT | GGGAAATAAG  | 3700 |
| AGGTTTCCAG | CCCAAAGAGA  | AGGAAAGACT | ATGTGGTGTT | ACTCTAAAAA  |      |
| GTATTTAATA | ACCGTTTTGT  | TGTTGCTGTT | GCTGTTTTGA | AATCAGATTG  | 3800 |
| TCTCCTCTCC | ATATTTTATT  | TACTTCATTC | TGTTAATTCC | TGTGGAATTA  |      |
| CTTAGAGCAA | GCATGGTGAA  | TTCTCAACTG | TAAAGCCAAA | TTTCTCCATC  | 3900 |
| ATTATAATTT | CACATTTTGC  | CTGGCAGGTT | ATAATTTTTA | TATTTCCACT  |      |
| GATAGTAATA | AGGTAAATATC | ATTACTTAGA | TGGATAGATC | TTTTTTCATAA | 4000 |
| AAAGTACCAT | CAGTTATAGA  | GGGAAGTCAT | GTTTCATGTT | AGGAAGGTCA  |      |
| TTAGATAAAG | CTTCTGAATA  | TATTATGAAA | CATTAGTTCT | GTCATTCTTA  | 4100 |
| GATTCTTTTT | GTTAAATAAC  | TTTAAAAGCT | AACTTACCTA | AAAGAAATAT  |      |
| CTGACACATA | TGAACCTCTC  | ATTAGGATGC | AGGAGAAGAC | CCAAGCCACA  | 4200 |
| GATATGTATC | TGAAGAATGA  | ACAAGATTCT | TAGGCCCGGC | ACGGTGGCTC  |      |
| ACATCTGTAA | TCTCAAGAGT  | TTGAGAGGTC | AAGGCGGGCA | GATCACCTGA  | 4300 |
| GGTCAGGAGT | TCAAGACCAG  | CCTGGCCAAC | ATGATGAAAC | CCTGCCTCTA  |      |
| CTAAAAATAC | AAAAATTAGC  | AGGGCATGGT | GGTGCATGCC | TGCAACCCTA  | 4400 |
| GCTACTCAGG | AGGCTGAGAC  | AGGAGAATCT | CTTGAACCTT | CGAGGCGGAG  |      |
| GTTGTGGTGA | GCTGAGATCC  | CTCTACTGCA | CTCCAGCCTG | GGTGACAGAG  | 4500 |
| ATGAGACTCC | GTCCCTGCCG  | CCGCCCCCGC | CTTCCCCCCC | AAAAAGATTTC |      |
| TTCTTCATGC | AGAACATACG  | GCAGTCAACA | AAGGGAGACC | TGGGTCCAGG  | 4600 |
| TGTCCAAGTC | ACTTATTTTCG | AGTAAATTAG | CAATGAAAGA | ATGCCATGGA  |      |
| ATCCCTGCCC | AAATACCTCT  | GCTTATGATA | TTGTAGAATT | TGATATAGAG  | 4700 |

FIGURE 1B

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|                 |             |            |            |             |      |
|-----------------|-------------|------------|------------|-------------|------|
| TTGTATCCCA      | TTTAAGGAGT  | AGGATGTAGT | AGGAAAGTAC | TAAAAACAAA  |      |
| CACACAAACA      | GAAAACCCTC  | TTTGCTTTGT | AAGGTGGTTC | CTAAGATAAT  | 4800 |
| GTCAGTGCAA      | TGCTGGAAT   | AATATTTAAT | ATGTGAAGGT | TTTAGGCTGT  |      |
| GTTTTCCCT       | CCTGTTCTTT  | TTTTCTGCCA | GCCCTTTGTC | ATTTTTGCAG  | 4900 |
| GTCAATGAAT      | CATGTAGAAA  | GAGACAGGAG | ATGAAACTAG | AACCAGTCCA  |      |
| TTTTGCCCT       | TTTTTTATTT  | TCTGGTTTTG | GTAAGAGATA | CAATGAGGTA  | 5000 |
| GGAGGTTGAG      | ATTTATAAAT  | GAAGTTTAAT | AAGTTTCTGT | AGCTTTGATT  |      |
| TTTCTCTTTC      | ATATTTGTTA  | TCTTGATAA  | GCCAGAAATG | GCCTGTAAAA  | 5100 |
| TCTACATATG      | GATATTGAAG  | TCTAAATCTG | TTCAACTAGC | TTACACTAGA  |      |
| TGGAGATATT      | TTCATATTCA  | GATACACTGG | AATGTATGAT | CTAGCCATGC  | 5200 |
| GTAATATAGT      | CAAGTGTGTT  | AAGGTATTTA | TTTTTAATAG | CGTCTTTAGT  |      |
| TGTGGACTGG      | TTCAAGTTTT  | TCTGCCAATG | ATTCTTCAA  | ATTTATCAAA  | 5300 |
| TATTTTTTCCA     | TCATGAAGTA  | AAATGCCCTT | GCAGTCACCC | TTCTTGAAGT  |      |
| TTGAACGACT      | CTGCTGTTTT  | AAACAGTTTA | AGCAAATGGT | ATATCATCTT  | 5400 |
| CCGTTTACTA      | TGTAGCTTAA  | CTGCAGGCTT | ACGCTTTTGA | GTCAGCGGCC  |      |
| AACTTTATTG      | CCACCTTCAA  | AAGTTTATTA | TAATGTTGTA | AATTTTACT   | 5500 |
| TCTCAAGGTT      | AGCATACTTA  | GGAGTTGCTT | CACAATTAGG | ATTCAGGAAA  |      |
| GAAAGAACTT      | CAGTAGGAAC  | TGATTGGAAT | TTAATGATGC | AGCATTCAT   | 5600 |
| GGGTACTAAT      | TTCAAAGAAT  | GATATTACAG | CAGACACACA | GCAGTTATCT  |      |
| TGATTTTCTA      | GGAATAATTG  | TATGAAGAAT | ATGGCTGACA | ACACGGCCTT  | 5700 |
| ACTGCCACTC      | AGCGGAGGCT  | GGACTAATGA | ACACCCTACC | CTTCTTTCCT  |      |
| TTCTCTCAC       | ATTTTCATGAG | CGTTTTGTAG | GTAACGAGAA | AATTGACTTG  | 5800 |
| CATTTGCATT      | ACAAGGAGGA  | GAACTGGCA  | AAGGGGATGA | TGGTGGAAGT  |      |
| TTTGTCTCTG      | CTAATGAAGT  | GAAAAATGAA | AATGCTAGAG | TTTTGTGCAA  | 5900 |
| CATAATAGTA      | GCAGTAAAAA  | CCAAGTGAAA | AGTCTTTCCA | AAACTGTGTT  |      |
| C               |             |            |            |             |      |
| AAGAGGGCAT      | CTGCTGGGAA  | ACGATTTGAG | GAGAAGGTAC | TAAATTGCTT  | 6000 |
| GGTATTTTCC      | GTAGGAACCC  | CAGAGCGAAA | TACAGTTTGC | AAAAGATGTC  |      |
| T               |             |            |            |             |      |
| [exon 3: 6015.. |             |            |            |             |      |
| CAGATGGGTT      | CTTCTCAAAT  | GAGACGTCAT | CTAAAGCACC | CTGTAGAAAA  | 6100 |
| CACACAAATT      | GCAGTGTCTT  | TGGTCTCCTG | CTAACTCAGA | AAGGAAATGC  |      |
| AACACACGAC      | AACATATGTT  | CCGGAACAG  | TGAATCAACT | CAAAAATGTG  | 6200 |
| GAATAGGTAA      | TTACATTCCA  | AAATACGTCT | TTGTACGATT | TTGTAGTATC  |      |
| ..6206]         |             |            |            |             |      |
| ATCTCTCTCT      | CTGAGTTGAA  | CACAAGGCCT | CCAGCCACAT | TCTTGGTCAA  | 6300 |
| ACTTACATTT      | TCCCTTTCTT  | GAATCTTAAC | CAGCTAAGGC | TACTCTCGAT  |      |
| GCATTACTGC      | TAAAGCTACC  | ACTCAGAATC | TCTCAAAAAC | TCATCTTCTC  | 6400 |
| ACAGATAACA      | CCTCAAAGCT  | TGATTTTCTC | TCCTTTCACA | CTGAAATCAA  |      |
| ATCTTGCCCA      | TAGGCAAAGG  | GCAGTGTCAA | GTTTGCCACT | GAGATGAAAT  | 6500 |
| TAGGAGAGTC      | CAAAGCTGAG  | AATTCACGTT | GTGTGTTATT | ACTTTCACGA  |      |
| ATGTCTGTAT      | TATTAATAA   | AGTATATATT | GGCAACTAAG | AAGCAAAGTG  | 6600 |
| ATATAACAT       | GATGACAAAT  | TAGGCCAGGC | ATGGTGGCTT | ACTCCTATAA  |      |
| TCCCAACATT      | TTGGGGGGCC  | AAGGTAGGCA | GATCACTTGA | GGTCAGGATT  | 6700 |
| TCAAGACCAG      | CCTGACCAAC  | ATGGTGAAAC | CTGTCTCTA  | CTAAAAATAC  |      |
| AAAAATTAGC      | TGGGCATGGT  | AGCAGGCAGT | TCTAGTACCA | GCTACTCAGG  | 6800 |
| GCTGAGGCAG      | GAGAATCGCT  | TGAACCCAGG | AGATGGAGGT | TGCAGTGAGC  |      |
| TGAGATTGTA      | CCACTGCACT  | CCAGTCTGGG | CAACAGAGCA | AGATTTTCATC | 6900 |
| ACACACACAC      | ACACACACAC  | ACACACACAC | ATTAGAAATG | TGTACTTGGC  |      |
| TTTGTTACCT      | ATGGTATTAG  | TGCATCTATT | GCATGGAAT  | TCCAAGCTAC  | 7000 |
| TCTGGTTGTG      | TTAAGCTCTT  | CATTGGGTAC | AGGTCACTAG | TATTAAGTTC  |      |
| AGGTTATTCG      | GATGCATTCC  | ACGGTAGTGA | TGACAATTCA | TCAGGCTAGT  | 7100 |
| GTGTGTGTTT      | ACCTTGTCAC  | TCCCACCACT | AGACTAATCT | CAGACCTTCA  |      |
| CTCAAAGACA      | CATTACACTA  | AAGATGATTT | GCTTTTTTGT | GTTTAATCAA  | 7200 |

FIGURE 1C

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|                 |             |             |             |            |      |
|-----------------|-------------|-------------|-------------|------------|------|
| GCAATGGTAT      | AAACCAGCTT  | GACTCTCCCC  | AAACAGTTTT  | TCGTACTACA |      |
| AAGAAGTTTA      | TGAAGCAGAG  | AAATGTGAAT  | TGATATATAT  | ATGAGATTCT | 7300 |
| AACCCAGTTC      | CAGCATTGTT  | TCATTGTGTA  | ATTGAAATCA  | TAGACAAGCC |      |
| ATTTTAGCCT      | TTGCTTTCTT  | ATCTAAAAAA  | AAAAAAAAAA  | AAATGAAGGA | 7400 |
| AGGGGTATTA      | AAAGGAGTGA  | TCAAATTTTA  | ACATTCTCTT  | TAATTAATTC |      |
| ATTTTAAATT      | TTACTTTTTT  | TCATTATTG   | TGCACTTACT  | ATGTGGTACT | 7500 |
| GTGCTATAGA      | GGCTTTAACA  | TTTATAAAAA  | CACTGTGAAA  | GTTGCTTCAG |      |
| ATGAATATAG      | GTAAGTAGAAC | GGCAGAACTA  | GTATTCAAAG  | CCAGGTCTGA | 7600 |
| TGAATCCAAA      | AACAAACACC  | CATTACTCCC  | ATTTTCTGGG  | ACATACTTAC |      |
| TCTACCCAGA      | TGCTCTGGGC  | TTTGTAATGC  | CTATGTAAAT  | AACATAGTTT | 7700 |
| TATGTTTGGT      | TATTTTCCTA  | TGTAATGTCT  | ACTTATATAT  | CTGTATCTAT |      |
| CTCTTGCTTT      | GTTTCCAAAG  | GTAAACTATG  | TGCTCTAAATG | TGGGCAAAAA | 7800 |
| ATAACACACT      | ATTCCAAATT  | ACTGTTCAAA  | TTCTTTTAAG  | TCAGTGATAA |      |
| TTATTTGTTT      | TGACATTAAT  | CATGAAGTTC  | CCTGTGGGTA  | CTAGGTAAAC | 7900 |
| CTTTAATAGA      | ATGTTAATGT  | TTGTATTTCAT | TATAAGAATT  | TTTGGCTGTT |      |
| ACTTATTTAC      | AACAATATTT  | CACTCTAATT  | AGACATTTAC  | TAAACTTTCT | 8000 |
| CTTGAAAACA      | ATGCCCAAAA  | AAGAACATTA  | GAAGACACGT  | AAGCTCAGTT |      |
| GGTCTCTGCC      | ACTAAGACCA  | GCCAACAGAA  | GCTTGATTTT  | ATTCAAACTT | 8100 |
| TGCATTTTAG      | CATATTTTAT  | CTTGGAAAAT  | TCAATTGTGT  | TGGTTTTTTG |      |
| A               |             |             |             |            |      |
| TTTTTGTTTG      | TATTGAATAG  | ACTCTCAGAA  | ATCCAATTGT  | TGAGTAAATC | 8200 |
| TTCTGGGTTT      | TCTAACCTTT  | CTTTAGATGT  | TACCCTGTGT  | GAGGAGGCAT |      |
| [exon 4: 8227.. |             |             |             |            |      |
| TCTTCAGGTT      | TGCTGTTTCT  | ACAAAGTTTA  | CGCCTAACTG  | GCTTAGTGTC | 8300 |
| TTGGTAGACA      | ATTTGCCTGG  | CACCAAAGTA  | AACGCAGAGA  | GTGTAGAGAG |      |
| T               |             |             |             |            |      |
| GATAAAACGG      | CAACACAGCT  | CACAAGAACA  | GACTTTCCAG  | CTGCTGAAGT | 8400 |
| G               |             |             |             |            |      |
| TATGGAAACA      | TCAAAACAAA  | GACCAAGATA  | TAGTCAAGAA  | GATCATCCAA |      |
| G               |             |             |             |            |      |
| GGTATGATAA      | TCTAAAATAA  | AAAGATCAAT  | CAGAAATCAA  | AGACACCTAT | 8500 |
| C               |             |             |             |            |      |
| ..8451]         |             |             |             |            |      |
| TTATCATAAA      | CCAGGAACAA  | GACTGCATGT  | ATGTTTAGTT  | GTGTGGATCT |      |
| TGTTTCCCTG      | TTGGAATCAT  | TGTTGGACTG  | AAAAAGTTTC  | CACCTGATAA | 8600 |
| TGTAGATGTG      | ATTCCACAAA  | CAGTTATACA  | AGGTTTTGTT  | CTCACCCCTG |      |
| CTCCCCAGTT      | TCCTTGTAAG  | GTATGTTGAA  | CACTCTAAGA  | GAAGAGAAAT | 8700 |
| GCATTTGAAG      | GCAGGGCTGT  | ATCTCAGGGA  | GTCGCTTCCA  | GATCCCTTAA |      |
| CGCTTCTGTA      | AGCAGCCCCCT | CTAGACCACC  | AAGGAGAAGC  | TCTATAACCA | 8800 |
| CTTTGTATCT      | TACATTGCAC  | CTCTACCAAG  | AAGCTCTGTT  | GTATTTACTT |      |
| GGTAATTCTC      | TCCAGGTAGG  | CTTTTCGTAG  | CTTACAAATA  | TGTTCTTATT | 8900 |
| AATCCTCATG      | ATATGGCCTG  | CATTAAAATT  | ATTTTAATGG  | CATATGTTAT |      |
| GAGAATTAAT      | GAGATAAAAT  | CTGAAAAGTG  | TTTGAGCCTC  | TTGTAGGAAA | 9000 |
| AAGCTAGTTA      | CAGCAAAATG  | TTCTCACATC  | TTATAAGTTT  | ATATAAAGAT |      |
| TCTCCTTTAG      | AAATGGTGTG  | AGAGAGAAAC  | AGAGAGAGAT  | AGGGAGAGAA | 9100 |
| GTGTGAAAGA      | ATCTGAAGAA  | AAGGAGTTTC  | ATCCAGTGTG  | GACTGTAAGC |      |
| TTTACGACAC      | ATGATGGAAA  | GAGTTCTGAC  | TTCAGTAAGC  | ATTGGGAGGA | 9200 |
| CATGCTAGAA      | GAAAAAGGAA  | GAAGAGTTTC  | CATAATGCAG  | ACAGGGTCAG |      |
| TGAGAAATTC      | ATTCAGGTCC  | TCACCAGTAG  | TTAAATGACT  | GTATAGTCTT | 9300 |
| GCACTACCTT      | AAAAAACTTC  | AAGTATCTGA  | AACCGGGGCA  | ACAGATTTTA |      |
| GGAGACCAAC      | GTCTTTGAGA  | GCTGATTGCT  | TTTGCTTATG  | CAAAGAGTAA | 9400 |
| ACTTTTATGT      | TTTGAGCAAA  | CCAAAAGTAT  | TCTTTGAACG  | TATAATTAGC |      |
| CCTGAAGCCG      | AAAGAAAAGA  | GAAAATCAGA  | GACCGTTAGA  | ATTGGAAGCA | 9500 |
| ACCAAATTCC      | CTATTTTATA  | AATGAGGACA  | TTTTAACCCA  | GAAAGATGAA |      |

FIGURE 1D

|                  |             |             |            |             |       |
|------------------|-------------|-------------|------------|-------------|-------|
| 5/7              |             |             |            |             |       |
| CCGATTTGGC       | TTAGGGCTCA  | CAGATACTAA  | GTGACTCATG | TCATTAATAG  | 9600  |
| AAATGTTAGT       | TCCTCCCTCT  | TAGGTTTGTA  | CCCTAGCTTA | TTACTGAAAT  |       |
| ATTCTCTAGG       | CTGTGTGTCT  | CCTTTAGTTC  | CTCGACCTCA | TGTCTTTGAG  | 9700  |
| TTTTTCAGATA      | TCCTCCTCAT  | GGAGGTAGTC  | CTCTGGTGCT | ATGTGTATTCT |       |
| TTTAAAGGCT       | AGTTACGGCA  | ATTAACCTTAT | CAACTAGCGC | CTACTAATGA  | 9800  |
| AACTTTGTAT       | TACAAAGTAG  | CTAACTTGAA  | TACTTTCCTT | TTTTTCTGAA  |       |
| ATGTTATGGT       | GGTAATTTCT  | CAAACCTTTTT | CTTAGAAAAC | TGAGAGTGAT  | 9900  |
| GTGTCTTATT       | TTCTACTGTT  | AATTTTCAAA  | ATTAGGAGCT | TCTTCCAAAG  |       |
| TTTTGTTGGA       | TGCCAAAAAT  | ATATAGCATA  | TTATCTTATT | ATAACAAAAA  | 10000 |
| ATATTTATCT       | CAGTTCTTAG  | AAATAAATGG  | TGTCACTTAA | CTCCCTCTCA  |       |
| AAAGAAAAGG       | TTATCATTGA  | AATATAATTA  | TGAAATTCTG | CAAGAACCTT  | 10100 |
| TTGCCTCAGC       | CTTGTTTTAT  | GATGGCATTG  | GATGAATATA | AATGATGTGA  |       |
| ACACTTATCT       | GGGCTTTTGC  | TTTATGCAGA  | TATTGACCTC | TGTGAAAACA  | 10200 |
| [exon 5: 10180.. |             |             |            |             |       |
| GCGTGCAGCG       | GCACATTGGA  | CATGCTAACC  | TCACCTTCGA | GCAGCTTCGT  |       |
| A                |             |             |            |             |       |
| AGCTTGATGG       | AAAGCTTACC  | GGGAAAGAAA  | GTGGGAGCAG | AAGACATTGA  | 10300 |
| AAAAACAATA       | AAGGCATGCA  | AACCCAGTGA  | CCAGATCCTG | AAGCTGCTCA  |       |
| GTTTGTGGCG       | AATAAAAAAT  | GGCGACCAAG  | ACACCTTGAA | GGGCCTAATG  | 10400 |
| CACGCACTAA       | AGCACTCAAA  | GACGTACCAC  | TTTCCCAAAA | CTGTCACTCA  |       |
| GAGTCTAAAG       | AAGACCATCA  | GGTTCCTTCA  | CAGCTTCACA | ATGTACAAAT  | 10500 |
| TGTATCAGAA       | GTTATTTTTA  | GAAATGATAG  | GTAACCAGGT | CCAATCAGTA  |       |
| C                |             |             |            |             |       |
| AAAATAAGCT       | GCTTATAACT  | GGAAATGGCC  | ATTGAGCTGT | TTCTCACAA   | 10600 |
| ..10568]         |             |             |            |             |       |
| TTGGCGAGAT       | CCCATGGATG  | AGTAACTGT   | TTCTCAGGCA | CTTGAGGCTT  |       |
| TCAGTGATAT       | CTTTCTCATT  | ACCACTGACT  | AATTTTGCCA | CAGGGTACTA  | 10700 |
| AAAGAAACTA       | TGATGTGGAG  | AAAGGACTAA  | CATCTCCTCC | AATAAACCCC  |       |
| AAATGGTTAA       | TCCAACCTGTC | AGATCTGGAT  | CGTTATCTAC | TGACTATATT  | 10800 |
| TTCCCTTATT       | ACTGCTTGCA  | GTAATTCAAC  | TGGAAATTAA | AAAAAAAAAA  |       |
| CTAGACTCCA       | CTGGGCCCTTA | CTAAATATGG  | GAATGTCTAA | CTTAAATAGC  | 10900 |
| TTTGGGATTC       | CAGCTATGCT  | AGAGGCTTTT  | ATTAGAAAGC | CATATTTTTT  |       |
| TCTGTAAAAG       | TTACTAATAT  | ATCTGTAACA  | CTATTACAGT | ATTGCTATTT  | 11000 |
| ATATTCATTC       | AGATATAAGA  | TTTGACATA   | TTATCATCCT | ATAAAGAAAC  |       |
| GGTATGACTT       | AATTTTAGAA  | AGAAAATTAT  | ATTCTGTTTA | TTATGACAAA  | 11100 |
| TGAAAGAGAA       | AATATATATT  | TTAATGGAA   | AGTTTGTAGC | ATTTTTCTAA  |       |
| TAGGTACTGC       | CATATTTTTT  | TGTGTGGAGT  | ATTTTTATAA | TTTTATCTGT  | 11200 |
| ATAAGCTGTA       | ATATCATTTT  | ATAGAAAATG  | CATTATTTAG | TCAATTGTTT  |       |
| AATGTTGGAA       | AACATATGAA  | ATATAAATTA  | TCTGAATATT | AGATGCTCTG  | 11300 |
| AGAAATTGAA       | TGTACCTTAT  | TTAAAAGATT  | TTATGGTTTT | ATAACTATAT  |       |
| AAATGACATT       | ATTAAAGTTT  | TCAAATTATT  | TTTTATTGCT | TTCTCTGTTG  | 11400 |
| CTTTTTATT        |             |             |            |             | 11408 |

FIGURE 1E

## POLYMORPHISMS IN THE CODING SEQUENCE OF TNFRSF11B

|             |            |            |             |            |      |
|-------------|------------|------------|-------------|------------|------|
| ATGAACAAGT  | TGCTGTGCTG | CGCGCTCGTG | TTTCTGGACA  | TCTCCATTAA |      |
| C           |            |            |             |            |      |
| GTGGACCACC  | CAGGAAACGT | TTCTCCAAA  | GTACCTTCAT  | TATGACGAAG | 100  |
| AAACCTCTCA  | TCAGCTGTTG | TGTGACAAAT | GTCCTCCTGG  | TACCTACCTA |      |
| AAACAACACT  | GTACAGCAAA | GTGGAAGACC | GTGTGCGCCC  | CTTGCCCTGA | 200  |
| CCACTACTAC  | ACAGACAGCT | GGCACACCAG | TGACGAGTGT  | CTATACTGCA |      |
| GCCCCGTGTG  | CAAGGAGCTG | CAGTACGTCA | AGCAGGAGTG  | CAATCGCACC | 300  |
| CACAACCGCG  | TGTGCGAATG | CAAGGAAGGG | CGCTACCTTG  | AGATAGAGTT |      |
| CTGCTTGAAA  | CATAGGAGCT | GCCCTCCTGG | ATTTGGAGTG  | GTGCAAGCTG | 400  |
| GAACCCGAGA  | GCGAAATACA | GTTTGCAAAA | GATGTCCAGA  | TGGGTTCTTC |      |
| TCAAATGAGA  | CGTCATCTAA | AGCACCTGT  | AGAAAACACA  | CAAATTGCAG | 500  |
| TGTCTTTGGT  | CTCCTGCTAA | CTCAGAAAGG | AAATGCAACA  | CACGACAACA |      |
| TATGTTCCGG  | AAACAGTGAA | TCAACTCAAA | AATGTGGAAT  | AGATGTTACC | 600  |
| CTGTGTGAGG  | AGGCATTCTT | CAGGTTTGCT | GTTCTTACAA  | AGTTTACGCC |      |
| TAACTGGCTT  | AGTGTCTTGG | TAGACAATTT | GCCTGGCACC  | AAAGTAAACG | 700  |
|             |            |            | T           |            |      |
| CAGAGAGTGT  | AGAGAGGATA | AAACGGCAAC | ACAGCTCACA  | AGAACAGACT |      |
|             | G          |            |             |            |      |
| TTCCAGCTGC  | TGAAGTTATG | GAAACATCAA | AACAAAGACC  | AAGATATAGT | 800  |
|             | G          |            |             |            |      |
| CAAGAAGATC  | ATCCAAGATA | TTGACCTCTG | TGAAAACAGC  | GTGCAGCGGC |      |
|             |            |            | A           |            |      |
| ACATTGGACA  | TGCTAACCTC | ACCTTCGAGC | AGCTTCGTAG  | CTTGATGGAA | 900  |
| AGCTTACCGG  | GAAAGAAAGT | GGGAGCAGAA | GACATTGAAA  | AAACAATAAA |      |
| GGCATGCAAA  | CCCAGTGACC | AGATCCTGAA | GCTGCTCAGT  | TTGTGGCGAA | 1000 |
| TAAAAAATGG  | CGACCAAGAC | ACCTTGAAGG | GCCTAATGCA  | CGCACTAAAG |      |
| CACTCAAAGA  | CGTACCACTT | TCCCAAAACT | GTCACCTCAGA | GTCTAAAGAA | 1100 |
| GACCATCAGG  | TTCTTTCACA | GCTTCACAAT | GTACAAATTG  | TATCAGAAGT |      |
|             |            |            | C           |            |      |
| TATTTTTTAGA | AATGATAGGT | AACCAGGTCC | AATCAGTAAA  | AATAAGCTGC | 1200 |
| TTATAA      |            |            |             |            | 1206 |

FIGURE 2

## ISOFORMS OF THE TNFRSF11B PROTEIN

|            |            |            |            |            |     |
|------------|------------|------------|------------|------------|-----|
| MNKLCCALV  | FLDISIKWTT | QETFPPKYLH | YDEETSHQLL | CDKCPPGTYL |     |
| N          |            |            |            |            |     |
| KQHCTAKWKT | VCAPCPDHYY | TDSWHTSDEC | LYCSPVCKEL | QYVKQECNRT | 100 |
| HNRVCECKEG | RYLEIEFCLK | HRSCPPGFGV | VQAGTPERNT | VCKRCPDGFF |     |
| SNETSSKAPC | RKHTNCSVFG | LLLTQKGNAT | HDNICSGNSE | STQKCGIDVT | 200 |
| LCEEAFRFA  | VPTKFTPNWL | SVLVDNLPGT | KVNAESVERI | KRQHSSQEQT |     |
|            |            |            | M          |            |     |
| FQLLKLWKHQ | NKDQDIVKKI | IQDIDLCENS | VQRHIGHANL | TFEQLRSLME | 300 |
|            |            |            | M          |            |     |
| SLPGKKVGAE | DIEKTIKACK | PSDQILKLLS | LWRIKNGDQD | TLKGLMHALK |     |
| HSKTYHFPKT | VTQSLKKTIR | FLHSFTMYKL | YQKLFLEMIG | NQVQSVKISC | 400 |
| L          |            |            |            |            | 401 |

10044574.010902

FIGURE 3



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 Stephens, J. Claiborne  
  
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Glu Glu Thr Ser His Gln Leu Leu Cys Asp Lys Cys Pro Pro Gly Thr
      35      40      45
Tyr Leu Lys Gln His Cys Thr Ala Lys Trp Lys Thr Val Cys Ala Pro
      50      55      60
Cys Pro Asp His Tyr Tyr Thr Asp Ser Trp His Thr Ser Asp Glu Cys
      65      70      75      80
Leu Tyr Cys Ser Pro Val Cys Lys Glu Leu Gln Tyr Val Lys Gln Glu
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Cys Asn Arg Thr His Asn Arg Val Cys Glu Cys Lys Glu Gly Arg Tyr
      100      105      110
Leu Glu Ile Glu Phe Cys Leu Lys His Arg Ser Cys Pro Pro Gly Phe
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Gly Val Val Gln Ala Gly Thr Pro Glu Arg Asn Thr Val Cys Lys Arg
      130      135      140
Cys Pro Asp Gly Phe Phe Ser Asn Glu Thr Ser Ser Lys Ala Pro Cys
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Arg Lys His Thr Asn Cys Ser Val Phe Gly Leu Leu Leu Thr Gln Lys
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      180      185      190
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Trp Lys His Gln Asn Lys Asp Gln Asp Ile Val Lys Lys Ile Ile Gln
      260      265      270
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Asn Leu Thr Phe Glu Gln Leu Arg Ser Leu Met Glu Ser Leu Pro Gly
      290      295      300
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206070-49404

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|     |     |     |     | 325 |     |     |     |     | 330 |     |     |     |     | 335 |     |  |  |  |  |
| Gly | Asp | Gln | Asp | Thr | Leu | Lys | Gly | Leu | Met | His | Ala | Leu | Lys | His | Ser |  |  |  |  |
|     |     |     | 340 |     |     |     |     | 345 |     |     |     |     | 350 |     |     |  |  |  |  |
| Lys | Thr | Tyr | His | Phe | Pro | Lys | Thr | Val | Thr | Gln | Ser | Leu | Lys | Lys | Thr |  |  |  |  |
|     |     | 355 |     |     |     |     | 360 |     |     |     |     | 365 |     |     |     |  |  |  |  |
| Ile | Arg | Phe | Leu | His | Ser | Phe | Thr | Met | Tyr | Lys | Leu | Tyr | Gln | Lys | Leu |  |  |  |  |
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| tatgtgtctg | gagtgcctcc  | aaaggaccay | tgctcagagg | aatactttgc | cactacaggg | 1140 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1200 |
| tgaaaatgct | agagttttgt  | gcaacataay | agtagcagta | aaaaccaagt | gaaaagtctt | 1260 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1320 |
| gagaaggtac | taaattgctt  | ggtattttcy | gtaggaacc  | cagagcgaaa | tacagtgtgc | 1380 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1440 |
| gcttgatttt | attcaaaact  | tgcattttar | catattttat | cttggaatat | tcaattgtgt | 1500 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1560 |
| gtagacaatt | tgcttggcac  | caaagtaaay | gcagagagt  | tagagaggat | aaaacggcaa | 1620 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1680 |
| aaagtaaacg | cagagagtgt  | agagaggatr | aaacggcaac | acagctcaca | agaacagact | 1740 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1800 |
| caagaacaga | ctttccagct  | gctgaagttr | tggaaacatc | aaaacaaaga | ccaagatata | 1860 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 1920 |
| atagtcaaga | agatcatcca  | aggtatgatm | atctaaaata | aaaagatcaa | tcagaaatca | 1980 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 2040 |
| atgcagatat | tgacctctgt  | gaaaacagcr | tcgagcggca | cattggacat | gctaacctca | 2100 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 2160 |
| gcttcacaat | gtacaaattg  | tatcagaagy | tatttttaga | aatgatagg  | aaccaggtcc | 2220 |
| nnnnnnnnnn | nnnnnnnnnn  | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | nnnnnnnnnn | 2280 |